STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
 U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

	10/60 7910
ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10652,791 B
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE	
IWrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters , instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s)contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
11Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12Patentin 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13Visuse of n/Xaa	"n" can only represent a single <u>nucleotide</u> ; "Xaa" can only represent a single <u>amino acid</u>
AMC – Biotechnology Systems Branch – 09/09/2003	



IFW16

RAW SEQUENCE LISTING

DATE: 10/03/2006

PATENT APPLICATION: US/10/652,791B

TIME: 10:37:42

Input Set: E:\03-332-3_Sep 2006.txt
Output Set: N:\CRF4\10032006\J652791B.raw

```
3 <110 > APPLICANT: Sirna Therapeutics, Inc.
             McSwiggen, James
     5
             Chowrira, Bharat
     7 <120> TITLE OF INVENTION: RNA Interference Mediated Inhibition of Platelet-Derived
             Endothelial Cell Growth Factor (ECGF1) Gene Expression Using
             Short Interfering Nucleic Acid (siNA)
    11 <130> FILE REFERENCE: 400/126 (MBHB 03-332-B)
    13 <140> CURRENT APPLICATION NUMBER: US 10/652,791B
    14 <141> CURRENT FILING DATE: 2003-08-29
    16 <150> PRIOR APPLICATION NUMBER: US 10/422,704
                                                               Does Not Comply
    17 <151> PRIOR FILING DATE: 2003-04-24
                                                               Corrected Diskette Needed
    19 <150> PRIOR APPLICATION NUMBER: US 10/417,012
    20 <151> PRIOR FILING DATE: 2003-04-16
    22 <150> PRIOR APPLICATION NUMBER: PCT/US 03/05346
    23 <151> PRIOR FILING DATE: 2003-02-20
    25 <150> PRIOR APPLICATION NUMBER: PCT/US 03/05028
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    28 <150> PRIOR APPLICATION NUMBER: US 60/358,580
    29 <151> PRIOR FILING DATE: 2002-02-20
    31 <150> PRIOR APPLICATION NUMBER: US 60/363,124
    32 <151> PRIOR FILING DATE: 2002-03-11
    34 <150> PRIOR APPLICATION NUMBER: US 60/386,782
    35 <151> PRIOR FILING DATE: 2002-06-06
    37 <150> PRIOR APPLICATION NUMBER: US 60/406,784
    38 <151> PRIOR FILING DATE: 2002-08-29
    40 <150> PRIOR APPLICATION NUMBER: US 60/408,378
    41 <151> PRIOR FILING DATE: 2002-09-05
    43 <150> PRIOR APPLICATION NUMBER: US 60/409,293
    44 <151> PRIOR FILING DATE: 2002-09-09
    46 <150> PRIOR APPLICATION NUMBER: US 60/440,129
    47 <151> PRIOR FILING DATE: 2003-01-15
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sense region
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DATE: 10/03/2006

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PATENT APPLICATION: US/10/652,791B
                                                            TIME: 10:37:42
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    74 ccqcugugcg cgaacccug
                                                           And Burker the Commence
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    78 <211> LENGTH: 19
    79 <212> TYPE: RNA
    80 <213> ORGANISM: Artificial Sequence
    82 <220> FEATURE:
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                                                      1. 5.
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     92 <213> ORGANISM: Artificial Sequence
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RAW SEQUENCE LISTING

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DATE: 10/03/2006 RAW SEQUENCE LISTING TIME: 10:37:42 PATENT APPLICATION: US/10/652,791B Input Set : E:\03-332-B Sep 2006.txt Output Set: N:\CRF4\10032006\J652791B.raw

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211 <212> TYPE: RNA

212 <213> ORGANISM: Artificial Sequence

RAW SEQUENCE LISTING DATE: 10/03/2006 PATENT APPLICATION: US/10/652,791B TIME: 10:37:42

Input Set : E:\03-332-B_Sep 2006.txt
Output Set: N:\CRF4\10032006\J652791B.raw

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RAW SEQUENCE LISTING DATE: 10/03/2006
PATENT APPLICATION: US/10/652,791B TIME: 10:37:42

Input Set : E:\03-332-B_Sep 2006.txt
Output Set: N:\CRF4\10032006\J652791B.raw

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356 <213> ORGANISM: Artificial Sequence

358 <220> FEATURE:

359 <223> OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense region

10/652,791B

Page 10

<210> 207 <211> 21 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: siNA sense region Unich Which pibonucleotive pibonucleotive does "N" represent.? represent.? <220> misc feature <221> <222> '(1)..(1) <223> 5'-3 attached terminal deoxyabasic moeity <220> <221> misc feature <222> (21)..(21) <223> 3'-3 attached terminal deoxyabasic moeity <220> <221> misc_feature <222> (1)..(19) n stands for any ribonucleotide <<u>4</u>00> 207 nnnnnnnnn nnnnnnnnt t See item#13 On error summary Theet. n typo of errors shown exist throughout the Sequence Listing. Please check subsequent

coquonicos for similar errors.

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/652,791B

DATE: 10/03/2006 TIME: 10:37:43

Input Set : E:\03-332-B_Sep 2006.txt
Output Set: N:\CRF4\10032006\J652791B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:207; N Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19
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Seq#:209; N Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19
Seq#:210; N Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19
Seq#:211; N Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19
Seq#:212; N Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19
Seq#:213; N Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19
Seq#:214; N Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19
Seq#:215; N Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19
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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

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Seq#:209; Line(s) 3349
Seq#:210; Line(s) 3371
Seq#:211; Line(s) 3398
Seq#:212; Line(s) 3425
Seq#:213; Line(s) 3452
Seq#:214; Line(s) 3480
Seq#:215; Line(s) 3507
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VERIFICATION SUMMARY

DATE: 10/03/2006 PATENT APPLICATION: US/10/652,791B TIME: 10:37:43

Input Set : E:\03-332-B_Sep 2006.txt

Output Set: N:\CRF4\10032006\J652791B.raw

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L:3308 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:207 after pos.:0
L:3335 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:208 after pos.:0
L:3357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:209 after pos.:0
L:3384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:210 after pos.:0
L:3411 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:211 after pos.:0
L:3438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:212 after pos.:0
L:3466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:213 after pos.:0
L:3493 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:214 after pos.:0
L:3521 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:215 after pos.:0
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